Q :	(AP	7AP 7AP 5AP 5AP	VAP AAP SAP SAP YGP/ AAP	TAP SAP AFP TSY TTY VTY VTY	FAP FAP FAP FAP FAP	
20	LOVHA	LDIV		2344445	\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\	
· 'C'	VGPE AGSE	VGAE YGAG YGAG YGPE		-YGPVSN -SCGQTA -FGTA -XCSS	-YGRC -YGRC -YGSC -YGSV -YGSV -YGSV -YGSV	
1	5.	55 50 50	~ !	YSE YSE YSE YSE	FSN FSN FSN FSN FSN FSN FSN FSN	
6.1 07.1	PA PA	PARYA- PARYA- PARYA- PARYS-		TY TY TY TY TY TY	SPARVA-SPARVT-SPARVT-SPASEP-ISPASEE-ISPASAR-ISPAR-IS	
,	TVGY	515Y 515Y 6VNY	SI SI SI SXKV	NDG NOC		
			HKKR (SL		
0	688 687 680	2 1 1 1 1 1 2 1 1 1 1 2	DERTE	YNIPEACK SVDQGDKCD QGDKCD Y-DDCN KENDSCN TEGDNCN TEGDSCN	ADACK	
9 7	S EGS S EGS S SGS		KN EGD NK AGN NL ENQ SR DAS AA TDL KN ESR KK DGI SK DSS	RIG COR NKG COR NKG GGR NKG GGR NKG GGR NTKG GGR NTEG GGR	VOS SNA- INA DNR IAA DNA QOS NNA QOS NNA TSA EAT YSR DNQ YSR DNQ YSR DNQ YSR DNQ	· · ·
1.40	AVDKAV VVOKAV AVDKAY	QAVURAY QAVNSAT QAVNQAT LAVNRAN DAVKKAY	EAVKNAV QAVNYAW RMIDKTT NAINGAV DMVNAWR KKVVNYAR KKVVNYAR KKVVNYAR	LLRA HDYA FEYG MADG FFRG FIEG	DSAVQSA DTAVMNA DNAVKNS NSAAANL NSAAANL NSAAANL NRAAAEI NNAVNTA NDAVENA DLAVNAA	
	A A A X X X X X X X X X X X X X X X X X	TAH ATL ATM STL KIL	VPEL -CSQNS -CSTTH -CSTTH -CDEWY FTVEAF VDYDAL NLPDET	TCGHSA PLPDST PLPDST GPGRLA GPRDVT GPRRLA GPRRLA GPCELA SAPDTLV	VAL TAL SSV SSV SAF SAF	
		1	MEI DE) - QPV 0) - KOK 0 - GKTV GKTV GKTV		
130	GPSG GPTG	GPSG SPSP SSAG STSG	STYGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	I APD-(PPDG-(PPDD PPDD PPDDD PPDDD	GGGGS GGGGYS GGGYS GGGYS GGGYS GGGYS GGGYS	*
	BASBPNG BSS168 G BSSDY G	BLSCARG BAPB92G BYSYABG BLS147G BSEPR G	BSISPLC TVTHER C DNEBPRC XCEXPRC BSBPF BSBPF EFCYLAC SPECPLAC	SHEXSP G AVPRCA G HMPPC3 G HSIPC2 G HSFURI G DHFURI G KLKEX1 G	VAPROA TAT11A TAPROK TAPROR TAPROT ACALPR AOALPR SCPRB1	*
)	. Section Page.	· • • • • • • • • • • • • • • • • • • •	· · · · · · · · · · · · · · · · · · ·	

• • • •

1

.

. ₹ · · · · •

